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Occupational exposure to bioburden in Portuguese bakeries: an approach to sampling viable microbial load

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In bakeries, a number of operations such as mixing are associated with exposure to air-suspended flour dust and related bioburden. The aim of this study was to find the best active sampling approach to the assessment of occupational exposure to bioburden in Portuguese bakeries based on the data obtained with the use of specific impaction and impinger devices. We used impaction to collect fungal particles from 100 L air samples onto malt extract agar (MEA) supplemented with chloramphenicol (0.05 %). For growing fungi we also used dichloran glycerol (DG18) agar-based media and for mesophilic bacteria we used tryptic soy agar (TSA) supplemented with nystatin (0.2 %). For *Enterobacteriaceae* we used violet red bile agar (VRBA). With impingers we also collected 300 L air samples at the 300 L/min airflow rate, inoculated onto the same culture media. The two methods, impaction and impinger, showed statistically significant differences in the following counts: fungal on MEA (z=-2.721, p=0.007), fungal on DG18 (z=-4.830, p=0.000), total bacteria (z=-5.435, p=0.000), and Gram-negative coliforms (z=-3.716, p=0.000). In all cases the impaction method detected significantly higher concentrations than the impinger method. Fungal and bacterial loads were higher in the production unit and lower in the shop. The fungal load obtained with impaction varied between 10 and 5140 CFU m⁻³, and total bacterial counts ranged between 10 and 4120 CFU m⁻³. This study has shown that the impaction method is the best active sampling approach to assessing viable bioburden in this specific occupational environment, but a multi-faceted approach to sampling and analyses combining methods and media enables a more refined risk characterisation and, consequently, better tailored risk control measures to reduce adverse health outcomes in workers.

KEY WORDS: bacteria; bioburden; fungi; impaction, impinger; malt extract agar; DG18 agar; tryptic soy agar; violet red bile agar

Exposure to bioaerosols and more specifically to the bioburden (microbial contaminants such as fungi and bacteria) is a critical occupational risk factor that requires deep understanding (1, 2). Some workplaces, such as agriculture, waste, fishery, forestry, mining, construction, and day care have already been associated with higher occupational exposure to biological hazards (1, 3). Some studies have reported higher prevalence of respiratory diseases and airway inflammation among workers in these sectors (4-7). Exposure to flour dust has been associated with asthma, conjunctivitis, rhinitis, and dermal reactions, while the so called baker's asthma has become one of the most severe and frequent expressions of occupational allergy (8).

Exposure to flour dust and related bioburden in bakeries mainly occurs during grinding, sifting, and mixing (8). When raw materials are mixed, profuse organic dust particles disseminate through the air and remain aerosolised for a long time before settling. Hence the high potential for exposure to bioburden and metabolites such as endotoxins, volatile organic compounds, and mycotoxins (9).

Sampling for bioburden in occupational environments can provide valuable information needed to characterise the exposure conditions, evaluate potential hazards to health, and establish control measures (10). Sampling methods are several and each situation is unique and requires specific methodology (3, 11). Theoretically, the most suitable way to assess indoor bioburden exposure is based on active air sampling. However, short sample times entail considerable drawbacks due to known large spatial and temporal variations (12). Bioburden has mostly been collected with stationary samplers, where exposure to microorganisms is calculated as time-weighted average of the airborne concentration at different sampling sites (10). Personal air sampling, in turn, provides a more accurate estimation of worker's exposure in the occupational environment (10). Epidemiological data from personal sampling have shown exposure-response associations almost twice as often as studies using stationary sampling (10, 13). Because of this, data obtained from stationary sampling need to be interpreted by a well-trained industrial hygienist following a more refined exposure assessment

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protocol with different sampling approaches or even different analytical methods.

Active bioburden sampling relies on three methods: impaction and impingement for stationary sampling and filtration for personal sampling (3, 14). Impactors use solid media such as agar to collect bioburden by impaction (3, 15). The number of colonies can be quantified by visual inspection after incubation, resulting in a direct viable bioburden count in a specific volume of air sampled (3, 15). This method is the recommended by the American Conference of Governmental Industrial Hygienists (ACGIH) (16) and by the Health Canada (17). Impinger sampling involves particle collection into liquid media. Normally, air is drawn by suction through a narrow inlet tube into a small flask containing the collection medium. Once sampling is complete, aliquots of the collection liquid can be cultivated in growth media to count viable microorganisms, since sample volumes and sampling times can be defined in advance (3, 15).

Although we can now rely on more refined analytical tools based on next-generation sequencing (NGS) technologies for occupational exposure assessments (18), culture-based methods will remain crucial, since they provide information about the infection potential of the bioburden present (19) and allow comparison with the existing guidelines (20).

The aim of this study was to find and discuss the best active sampling approach to assessing bioburden occupational exposure in bakeries based on data collected from impaction and impinger devices.

MATERIALS AND METHODS

As part of a larger exploratory study which received financial support from the Portuguese Authority for Working Conditions this study took place between May 2016 and June 2017 and included 14 Portuguese bakeries in the Lisbon district. Eight bakeries belonged to a company which also baked for school canteens and vending machines. Five were integrated in supermarkets owned by one owner. One produced only fresh dough and belonged to the same supermarket owner (Table 1).

Most bakeries had three working areas: production, where kneading machines and ovens were located and where dough was shaped; raw material warehouse, where workers stored raw materials for dough; and the shop, where baked bread and pastry were sold. Sampling sites and sampling periods were determined based on the high amount of time spent by workers on those places or dislocation frequency during their occupational activity (Table 1).

Bioburden sampling

We took 59 indoor air samples (100 L of air) using an impaction method at a flow rate of 140 L min⁻¹ (Millipore Air Tester, Millipore, Billerica, MA, USA), and the particles

were collected onto agar plates according to the manufacturer's instructions. We also took one outdoor sample for each bakery for reference (control). All indoor samples were collected from workplaces occupied by one or two workers. Three culture media were used for better selectivity between fungal and bacterial populations: malt extract agar (MEA) (Frilabo, Maia, Portugal) supplemented with chloramphenicol (0.05 %) and dichloran glycerol (DG18) (Frilabo) agar for fungi, tryptic soy agar (TSA) (Frilabo) supplemented with nystatin (0.2 %) for mesophilic bacteria, and violet red bile agar (VRBA) (Frilabo) for the *Enterobacteriaceae* (e.g. Gram-negative coliforms).

We also collected 59 air samples (300 L) using a Coriolis μ impinger (Bertin Technologies, Montigny-le-Bretonneux, France) at the airflow rate of 300 L min⁻¹. Samples were collected into vials with 10 mL sterile phosphate-buffered saline (pH 7.4) with 0.05 % Triton X. From each vial 150 μ L of the collection liquid was inoculated onto four culture media: 2 % MEA with 0.05 g L⁻¹ chloramphenicol media, DG18 agar, TSA with 0.2 % nystatin, and VRBA.

All samples were incubated at 27 °C for 5-7 days (fungi) or at 30 °C and 35 °C for 7 days (mesophilic bacteria and Gram-negative coliforms, respectively). After laboratory processing and incubation, the fungi and bacteria were counted and the counts expressed as colony-forming units per cubic metre of air (CFU m⁻³).

Statistical analysis

The data were analysed with the SPSS statistical software for Windows, v. 24.0 (IBM, Lisbon, Portugal). The threshold of significance was set at 5 % (p<0.05).

Table 1 Number of sampling sites in each bakery

Baker	ies	Indoor air sampling impaction (n)	Indoor air sampling impinger (n)
1	Company bakery 1	3+	3
2	Company bakery 2	5	5
3	Company bakery 3	4+	4+
4	Company bakery 4	4	4
5	Company bakery 5	5	5+
6	Company bakery 6	4+	4
7	Company bakery 7	5+	5+
8	Company bakery 8	4	4+
9	Supermarket bakery 1	4	4
10	Supermarket bakery 2	4	4+
11	Supermarket bakery 3	4	4
12	Supermarket bakery 4	3	3
13	Supermarket bakery 5	4	4
14	Fresh dough unit	6	6+
Total	ll the four media were used	59	59

	Minimum (CFU m ⁻³)			Interquartile Range 25–75 (CFU m ⁻³)	J	
Impaction Fungal Load MEA	10.0	5140.0	235.0	110.0-1210.0	7	
Impinger Fungal Load MEA	3.0	2620.0	85.0	13.0-310.0	18	
Impaction Fungal Load DG18	3.0	10310.0	140.0	17.0-740.0	9	
Impinger Fungal Load DG18	3.0	1670.0	10.0	7.0-80.0	33	
Impaction Total Bacteria Load TSA	10.0	4120.0	235.0	150.0-475.0	1	
Impinger Total Bacteria Load TSA	3.0	5306.7	7.0	3.3-28.5	20	
Impaction Gram- Bacte- ria Load VRB	4.0	50.0	10.0	10.0-10.0	53	
Impinger Gram-Bacteria Load VRB	3.0	3.3	3.2	3.0-3.3	69	

Table 2 Fungal and bacterial loads by method

Table 3 Comparison of the fungal and bacterial counts between the impaction and impinger method

			Ranks		Test statistics ^m		
		Ν	Mean rank	Sum of ranks	Z	р	
	Negative ranks	45ª	34.99	1574.50			
Impinger fungal load MEA –	Positive ranks	22 ^b	31.98	703.50	-2.721 ⁿ	0.007*	
Impaction fungal load MEA	Ties	1°			-2./21"	0.007*	
	Total 68						
	Negative ranks	47 ^d	33.43	1571.00		0.000*	
Impinger fungal load DG18 –	Positive ranks	13°	19.92	259.00	-4.830 ⁿ		
Impaction fungal load DG18	Ties	8 ^f			-4.850"		
	Total	68					
	Negative ranks	55 ^g	29.18	1605.00		0.000*	
Impinger total bacterial load -	Positive ranks	4 ^h	41.25	165.00	-5.435 ⁿ		
Impaction total bacterial load	Ties	2^i			-3.435		
	Total	61					
	Negative ranks	17 ^j	13.00	221.00		0.000*	
Impinger Gram-negative	Positive ranks	4 ^k	2.50	10.00	-3.716 ⁿ		
bacterial load – Impaction Gram-negative bacterial load					-3./10"	0.000	
5	Total	70					

a – impinger fungal load MEA<impaction fungal load MEA

b – *impinger fungal load MEA*>*impaction fungal load MEA*

c – *impinger fungal load MEA*=*impaction fungal load MEA*

d - impinger fungal load DG18<impaction fungal load DG18

 $e-impinger\ fungal\ load\ DG18{>}impaction\ fungal\ load\ DG18$

f – impinger fungal load DG18=impaction fungal load DG18

g – impinger total bacteria load<impaction total bacteria load

h – impinger total bacteria load>impaction total bacteria load

i – *impinger total bacteria load=impaction total bacteria load*

j – impinger coliform load<impaction coliform load *k* – impinger coliform load>impaction coliform load

l – impinger coliform load – impaction coliform load l – impinger coliform load – impaction coliform load

m - Wilcoxon signed ranks test

n – based on positive ranks

* statistically significant (p < 0.05)

Normality of distribution was tested with the Shapiro-Wilk test, but as it did not show normal distribution, we used the Wilcoxon test to compare the impaction and impinger methods for assessing fungal and bacterial loads.

We also compared the sensitivity and specificity between the impinger and impaction method.

For the comparison of fungal and bacterial loads (obtained with both methods) between the work areas (production, warehouse, and shop) we used the Kruskal-Wallis test.

RESULTS

Table 2 shows the fungal and bacterial loads by method. The two methods significantly differed in the following counts: fungal on MEA (z=-2.721, p=0.007), fungal on DG18 agar (z=-4.830, p=0.000), total bacteria (z=-5.435, p=0.000), and Gram-negative coliforms (z=-3.716, p=0.000). In all of these cases, impaction yielded significantly higher counts than the impinger method (Table 3).

Table 4 compares the sensitivity and specificity of the impaction and impinger methods for fungal loads on MEA and DG18 and bacterial loads on TSA and VRB. The impinger method was more sensitive for fungal loads on MEA (70.49 %) and on DG18 (53.33 %) and also for bacterial load on TSA (72.13 %). However, it was not nearly as specific (0.00 %, 15.15 %, and 5.56 %, respectively). In contrast, it was specific for Gram-negative coliforms (74.24 %) but not sensitive (0.00 %). The agreement between the two methods was also relatively low.

The three workplaces (production, warehouse, and the shop) did not significantly differ in either the fungal or bacterial load. However, the production area had almost consistently the highest fungal and bacterial loads (Figures 1 and 2).

DISCUSSION

Quite expectedly, the production area had the highest fungal and bacterial loads. Beside having more workers and raw materials out in the open (21, 22), this workplace has been reported for the highest particulate matter concentrations (23, 24), and particles are known bioburden carriers (25, 26).

However, the focus of this study was on the sampling methods. The impinger method is often used in occupational settings with higher microbial loads. Not only does this method allow dilution before incubation on plates but it also makes possible the use of molecular tools, as it produces liquid samples (3). This is not possible with samplers that employ impaction on solid media. On the downside, the impinger method yielded lower fungal counts than impaction, even though dry collection methods, such as impaction, usually report lower counts for stress-sensitive bioaerosols (27). The reason may be sought in liquid evaporation associated with longer operation with impingers, as it hampers fungal and bacterial viability (28).

Beside the sampling methods, the use of different fungal media (MEA and DG18 agar) allowed a more comprehensive insight into the problem, as it yielded different counts with the same sampling method, which has already been reported by others (20, 24). Yet both media showed good sensitivity to fungi, and the combination of the two media has been shown to give complementary information (29-31). As for the bacteria, TSA provides information about bacterial contamination and VRBA is more specific for the Gramnegative coliforms.

There are other factors, beside collection and media used, that can affect sampling, such as inlet efficiency, transmission efficiency, and collection efficiency (2, 10). Furthermore, the viable bioburden collected with both methods constitutes a small percentage of the total microbial

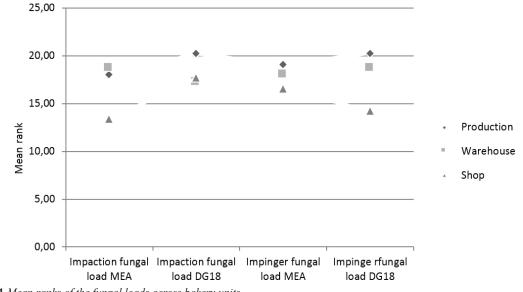


Figure 1 Mean ranks of the fungal loads across bakery units

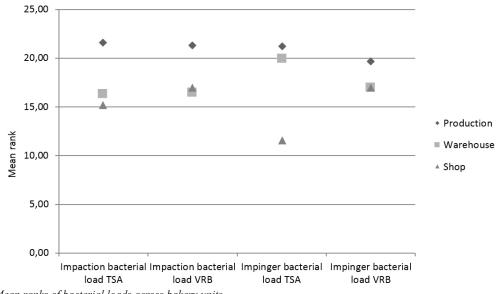


Figure 2 Mean ranks of bacterial loads across bakery units

Table 4 Sensitivity and	d specificity of the	impinger and	impaction method
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		Impinger fungal load MEA		Total	Sensitivity	Specificity	Agreement between
		Not detected	Detected			- F	the methods (%)
Impaction	Not detected	0	7	7	_		
fungal load MEA	Detected	18	43	61	70.49 %	0.09 %	63.24 %
Т	otal	18	50	68	-		
		Impinger fungal load					
		DG18		Total			
		Not detected	Detected		_		
Impaction	Not detected	5	3	8	53.33 %	15.15 %	54.41 %
fungal load DG18	Detected	28	32	60	-		
T	otal	33	35	68	-		
		Impinger total bacterial load		Total			
		Not detected	Detected				
Impaction	Not detected	1	0	1	72 12 0/	5 56 0/	72 58 0/
total bacte- ria load	Detected	17	44	61	72.13 %	5.56 %	72.58 %
Т	otal	18	44	62	-		
		Impinger Gram-nega-					
		tive bacterial load		Total			
		Not detected	Detected				
Impaction	Not detected	49	4	53	-		
Gram-neg- ative bacterial load	Detected	17	0	17	0.00 %	74.24 %	70.04 %
Total		66	4	70	-		

load (32) and this bias should be taken into consideration to properly interpret the obtained results (2, 20). The ability of a given airborne microbial population to grow on a medium is greatly affected by the physiological and physical stress caused by the aerosolisation, sampling methods, and other factors that affect airborne microbial cells (33, 34). Culture based-methods not only affect the microbial loads but also their biodiversity (35). In contrast, quantitative polymerase chain reaction (qPCR)-based methods can not differentiate between live and dead microorganisms, which is not totally suitable for infectious microorganisms and mainly occupational settings (36). Therefore, cultureindependent methods do not overcome all the limitations of the culture-based ones. While molecular tools such as next-generation sequencing (NGS) generally show greater biodiversity in aerosol samples than culture-based methods, more and more researchers recognise that culture-based and culture-independent methods are complementary (20, 36).

Given the complementarity of the culture basedmethods and molecular tools, we propose the following procedure: apply the impaction method to obtain information about the viable microbial load and the impinger method to target for specific microorganisms by later using molecular tools. This combination has already provided more comprehensive information about occupational exposure to bioburden (31, 37).

CONCLUSIONS

This study suggests that the impaction method is the best active sampling approach for occupational exposure assessment to viable bioburden in this specific occupational environment. Combining two or more fungal and/or bacterial media can provide even more information for exposure assessment. A multi-faceted approach to sampling and analyses is therefore welcome, as it enables a more refined risk characterisation and, consequently, better tailored risk control measures to reduce adverse health outcomes in workers.

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Conflict of interest

None.

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Pristup uzorkovanju mikroorganizama radi utvrđivanja profesionalnog biološkog opterećenja u portugalskim pekarama

Mnoge su djelatnosti u pekarama povezane s izloženosti lebdećim česticama brašna i pripadajućim mikroorganizmima, poput miješanja tijesta. Čilj je ovog istraživanja bio pronaći najbolji način aktivnog uzorkovanja radi ocjene profesionalne izloženosti biološkom opterećenju u portugalskim pekarama na temelju podataka dobivenih primjenom specifičnih impaktora i impingera. Impaktorom smo prikupljali čestice mikroskopski sitnih gljiva iz 100-litarskog uzorka zraka na agar od sladnog ekstrakta (engl. malt extract agar, krat. MAE) kojem je dodan kloramfenikol (0,05 %). Za rast gljiva također smo rabili dikloran glicerol (DG18) medij, a za mezofilne bakterije triptični sojin agar (TSA) s nistatinom (0,2 %). Za enterobakterije smo rabili ljubičasto-crveni žučni agar (engl. violet red bile agar, krat. VRBA). Impingerima smo također prikupljali uzorke zraka (300 L) uz protok od 300 L po minuti te ih cijepili na iste medije. Dvije su se metode, impakcijska i impingerska, statistički značajno razlikovale u sljedećim parametrima: u opterećenju gljivama na mediju MEA (z=-2,721, p=0,007), na mediju DG18 (z=-4,830, p=0,000), ukupnom bakterijskom opterećenju (z=-5,435, p=0,000) te u opterećenju gram-negativnim koliformima (z=-3,716, p=0,000). U svim je tim slučajevima impakcijska metoda dala značajno više koncentracije od impingerske. Opterećenje gljivama i bakterijama bilo je, sasvim očekivano, više u prostoriji za pripremu tijesta, a niže u prodavaonici. Opterećenje gljivama dobiveno impakcijom kretalo se u rasponu od 10 do 5140 CFU m⁻³, a bakterijama u rasponu od 10 do 4120 CFU m⁻³. Ovo je istraživanje izdvojilo impakcijsku metodu kao najbolji pristup aktivnom prikupljanju uzoraka za ocjenu opterećenja živim mikroorganizmima za ovaj specifični radni okoliš, ali bi višestrani pristup uzorkovanju i analizi, koji bi objedinjavao više metoda i medija, omogućio finiju karakterizaciju rizika, a time i mjere kontrole rizika koje bi bile bolje prilagođene specifičnim potrebama radi smanjenja štetnih zdravstvenih ishoda u radnika.

KLJUČNE RIJEČI: bakterije; biološko opterećenje; DG18 agar; gljive; impakcija, impinger; ljubičasto-crveni žučni agar; sladni agar; triptični sojni agar